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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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3945
1 MKKTRFPNKLNTLNTQRVLS......IAGISLGIWGIHTIRIRKHD 757
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1: sp_archea:*
2: sp_bacteria:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Result	Score	Query Match	% Query Match Length DB	DB	ID	1 1 1 1 1 1 1 1 1	Description
1	3941	99.9	757	2	Q8GRA2		O8gra2 streptococc
2	2058.5	52.2	756	N	Q8RP53		Q8rp53 streptococc
ω	1965.5	49.8	742	2	Q9ZB47		09zb47 streptococc
4	1862	47.2	744	16	Q8K8U7		08k8u7 streptococc
5	1254.5	31.8	524	16	Q8P2W3		08p2w3 streptococc
6	633.5	16.1	659	N	Q54953		054953 streptococc
7	633	16.0	869	N	Q8RP54		
8	611	15.5	685	N	033709		033709 streptococc
9	603	15.3	638	N	Q01924		
10	523	13.3	580	Ŋ	Q47942		
11	272.5	6.9	106	N	033715		
12	262.5	6.7	104	N	033714		033714 streptococc
13	241.5	6.1	108	N	033711		
14	236.5	6.0	340	16	'Q9A1S2		09als2 streptoco
15	224.5	5.7	696	16	Q8P2V7		08p2v7 streptococo
16	223	5.7	733	N	086961	•	Q8g9g1 streptococc

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## ALIGNMENTS

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121 SSVKKWYKHDGISTKFEDYAISPRITGDELNQKLRAVMYNGHPQNANGIMEGLEPLNAI 1	H	61 AINPDSSSEYRWYGYESYVRGHPYYKQFRVAHDLRVNLEGSRSYQVYCFNLKKAFPLGSD 1	٠. ب	1 MKKTRFPNKLNTLNTQRVLSKNSKRFTVTLVGVFLMIFALVTSMVGAKTVFGLVESSTPN 60	1 MKKTRFPNKLNTLNTQRVLSKNSKRFTVTLVGVFLMIFALVTSMVGAKTVFGLVESSTPN 60	Query Match 99.9%; Score 3941; DB 2; Length 757; Best Local Similarity 99.9%; Pred. No. 1.1e-209; Matches 756; Conservative 1; Mismatches 0; Indels 0; Gaps	SEQUENCE 757 AA; 85142 MW; 363414E6FF078DFD CRC64;	EMBL; AB083107; BAC20340.1;	Submitted (APR-2002) to the EMBL/GenBank/DDBJ databases.	The state of the s	*Characterizaition of Protein F-nomologous protein of M1-type group A	STRALIVACATION OF THE CONTROL OF THE	SEQUENCE FROM N.A.	[1]	NCBI_TaxID-1314;		Streptococous pyoguas; Bacteria; Firmicutes; Lactobacillales; Streptococcaceae;	Protein F-homologous protein.	(TrEMBLrel. 23, Last	(TrEMBLrel. 23,	Olymphr-2003 (AremBirel 23, Created)	Q8GRA2 PRELIMINARY; PRT; 757 AA.	RA2	
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NCBI_TaxID=1314;
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Bacteria; Firmicutes; Lactobacillales;
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Q9ZB47;
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NCBI_TaxID=1314;
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